## SEQUENCE LISTING

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<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapiens

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Ile Val Val Asn Glu Phe Cys Glu Arg Phe Ser Tyr Tyr Gly Met Lys 50 55 60

Ala Val Leu Ile Leu Tyr Phe Leu Tyr Phe Leu His Trp Asn Glu Asp 65 70 75 80

Thr Ser Thr Ser Ile Tyr His Ala Phe Ser Ser Leu Cys Tyr Phe Thr 85 90 95

Pro Ile Leu Gly Ala Ala Ile Ala Asp Ser Trp Leu Gly Lys Phe Lys
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Thr Ile Ile Tyr Leu Ser Leu Val Tyr Val Leu Gly His Val Ile Lys
115 120 125

Ser Leu Gly Ala Leu Pro Ile Leu Gly Gly Gln Val Val His Thr Val 130 \$135\$ 140

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Lys Pro Cys Val Ala Ala Phe Gly Gly Asp Gln Phe Glu Glu Lys His 165 170 175

Ala Glu Glu Arg Thr Arg Tyr Phe Ser Val Phe Tyr Leu Ser Ile Asn 180 185 190

Ala Gly Ser Leu Ile Ser Thr Phe Ile Thr Pro Met Leu Arg Gly Asp 195 200 205

Val Gln Cys Phe Gly Glu Asp Cys Tyr Ala Leu Ala Phe Gly Val Pro 210 215 220

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Lys	Cys	Ile	Trp 260	Phe	Ala	Ile	Ser	Asn 265	Arg	Phe	Lys	Asn	Arg 27.0	Ser	Gly
Asp	Ile	Pro 275	Lys	Arg	Gln	His	Trp 280	Leu	Asp	Trp	Ala	Ala 285	Glu	Lys	Tyr
Pro	Lys 290	Gln	Leu	Ile	Met	Asp 295	Val	Lys	Ala	Leu	Thr 300	Arg	Val	Leu	Phe
Leu 305	Tyr	Ile	Pro	Leu	Pro 310	Met	Phe	Trp	Ala	Leu 315	Leu	Asp	Gln	Gln	Gly 320
Ser	Arg	Trp	Thr	Leu 325	Gln	Ala	Ile	Arg	Met 330	Asn	Arg	Asn	Leu	Gly 335	Phe
Phe	Val	Leu	Gln 340	Pro	Asp	Gln	Met	Gln 345	Val	Leu	Asn	Pro	Phe 350	Leu	Val
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Ile	Asn	Glu	Met	Ala 405	Pro	Ala	Gln	Ser	Gly 410	Pro	Gln	Glu	Val	Phe 415	Leu
Gln	Val	Leu	Asn 420	Leu	Ala	Asp	Asp	Glu 425	Val	Lys	Val	Thr	Val 430	Val	Gly
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Thr Pro His Tyr Ser Lys Leu His Leu Lys Thr Lys Ser Gln Asp Phe His Phe His Leu Lys Tyr His Asn Leu Ser Leu Tyr Thr Glu His Ser Val Gln Glu Lys Asn Trp Tyr Ser Leu Val Ile Arg Glu Asp Gly Asn Ser Ile Ser Ser Met Met Val Lys Asp Thr Glu Ser Lys Thr Thr Asn Gly Met Thr Thr Val Arg Phe Val Asn Thr Leu His Lys Asp Val Asn Ile Ser Leu Ser Thr Asp Thr Ser Leu Asn Val Gly Glu Asp Tyr Gly Val Ser Ala Tyr Arg Thr Val Gln Arg Gly Glu Tyr Pro Ala Val His Cys Arg Thr Glu Asp Lys Asn Phe Ser Leu Asn Leu Gly Leu Leu Asp Phe Gly Ala Ala Tyr Leu Phe Val Ile Thr Asn Asn Thr Asn Gln Gly Leu Gln Ala Trp Lys Ile Glu Asp Ile Pro Ala Asn Lys Met Ser Ile . 605 Ala Trp Gln Leu Pro Gln Tyr Ala Leu Val Thr Ala Gly Glu Val Met Phe Ser Val Thr Gly Leu Glu Phe Ser Tyr Ser Gln Ala Pro Ser Ser Met Lys Ser Val Leu Gln Ala Ala Trp Leu Leu Thr Ile Ala Val Gly Asn Ile Ile Val Leu Val Val Ala Gln Phe Ser Gly Leu Val Gln Trp 

Ala Glu Phe Ile Leu Phe Ser Cys Leu Leu Leu Val Ile Cys Leu Ile 675 680 685 Phe Ser Ile Met Gly Tyr Tyr Tyr Val Pro Val Lys Thr Glu Asp Met Arg Gly Pro Ala Asp Lys His Ile Pro His Ile Gln Gly Asn Met Ile 710 715 Lys Leu Glu Thr Lys Lys Thr Lys Leu 725 <210> 3 <211> 363 DNA <212> <213> Homo Sapiens <400> gaggtccagc tgcaacagtc tggacctgag ctggtgaagc ctggagcttc aatgaagata 60 tcctgcaagg cttctggtta ctcattcact ggctacacca tgaactgggt gaagcagagc 120 catggaaaga accttgagtg gattggactt attaatcctt acaatggtgg tattaactac 180 aaccagaagt tcaagggcaa ggccacatta actgtagaca agtcatccag tacaqcctac 240 atggagetee teagtetgae atetgaggae tetgeagtet attactgtae aagaegggee 300 tactatggta actacggtac tatggactac tggggtcaag gaacctcagt caccgtctcc 360 tca 363 <210> <211> 321 <212> DNA <213> Homo Sapiens <400> 4 gaaaatgttc tcacccagtc tccagcaagc atgtctgcat ctccagggga aaaggtcacc 60 atgacctgca gtgccagctc aagtgtaagt tacatgcact ggtaccagca gaagtcaacc 120 acctcccca aactctggat ttatgacaca tccaatctgg cttctggggt cccaggtcgc 180 ttcagtggca gtgggtctgg aaactcttac tctctcacga tcagcaacat ggaggctgaa 240 gatgttgcca cttattactg ttttcagggg agtggttacc cactcacgtt cggtgctggg 300 accaagctgg agctgaaacg g 321

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cctggacaag	gccttgaatg	gattggtatg	attgatcctt	cagacagtga	aactcactac	180
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Ser Met Ly	s Ile Ser C		Ser Gly Tyr 25	Ser Phe Thr	Gly Tyr	
Thr Met As		ys Gln Ser 40	His Gly Lys	Asn Leu Glu 45	Trp Ile	

Gly Leu Ile Asn Pro Tyr Asn Gly Gly Ile Asn Tyr Asn Gln Lys Phe

50 55 60

Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr 65 70 75 80

Met Glu Leu Leu Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys 85 90 95

Thr Arg Arg Ala Tyr Tyr Gly Asn Tyr Gly Thr Met Asp Tyr Trp Gly 100 105 110

Gln Gly Thr Ser Val Thr Val Ser Ser 115 120

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<212> PRT

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Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser Ser Val Ser Tyr Met 20 25 30

His Trp Tyr Gln Gln Lys Ser Thr Thr Ser Pro Lys Leu Trp Ile Tyr 35 40 45

Asp Thr Ser Asn Leu Ala Ser Gly Val Pro Gly Arg Phe Ser Gly Ser 50 55 60

Gly Ser Gly Asn Ser Tyr Ser Leu Thr Ile Ser Asn Met Glu Ala Glu 65 70 75 80

Asp Val Ala Thr Tyr Cys Phe Gln Gly Ser Gly Tyr Pro Leu Thr
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Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg 100 105

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Trp Leu Asn Trp Val Arg Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile 35 40 45

Gly Met Ile Asp Pro Ser Asp Ser Glu Thr His Tyr Asn Gln Met Phe 50 55 60

Lys Asp Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr 65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys 85 90 95

Thr Ser Gln Gly Val Pro Val Pro Phe Asp Tyr Trp Gly Gln Gly Thr 100 105 110

Thr Leu Thr Val Ser Ser 115

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Asn Gly Asn Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser 35 40 45

Pro Lys Leu Leu Ile Tyr Arg Val Ser Asn Arg Phe Ser Gly Val Pro 50 55 60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile 65 70 75 80

Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser 85 90 95

Thr His Val Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys 100 105 110

Arg